

KCAGHLGYAADRNWKSKEWPDFVNSYASWWSSHVLDWLKYGKRLLVVHYEELRRSLVP TLREMVAFLNVSVSEERLLCVENNKEGSFRRRGRRSHDPEPFTPEMKDLINGYIRTVD QALRDHNWTGLPREYVPR"

BASE COUNT 1191 a 1340 c 1492 g 1149 t ORIGIN

1 gagetgegte agttgegteg eegetggtte caccacttea tgagtgacte eeagggaceg 61 cccgccetgg gccccgaggc tgccaggccc gccatccaca gccgaggcac ctacattgga 121 tgcttcagtg acgatggcca tgagaggact ctgaaaggag ctgtgtttta tgacttgaga 181 aagatgactg teteceactg ecaggatgeg tgtgetgage ggteetatgt etaegeegge 241 ttggaggccg gggcggagtg ttactgcggg aaccggctgc cagcggtgag cgtggggctg 301 gaagagtgta accatgagtg caaaggcgag aagggctctg tgtgcggggc tgtggaccgg 361 ctctccgtgt accgtgtgga cgagctgcag ccgggctcca ggaagcggcg gaccgccacc 421 taccgcggat gcttccgact gccagagaac atcacacatg ccttccccag ctccctgata 481 caggccaatg tgaccgtggg gacttgctcg ggcttttgtt cccagaaaga gttccccttg 541 gccattctca ggggctggga atgctactgt gcttacccta ccccccggtt caacctgcgg 601 gatgccatgg acagctcagt atgtggccag gaccctgagg cacagaggct ggcagaatac 661 tgtgaggtct accagacacc tgtgcaagac actcgttgta cagacaggag gttcctgcct 721 aacaaatcca aagtgtttgt ggctttgtca agcttcccag gagccgggaa cacgtgggca 781 cggcacctca ttgagcatgc cactggcttc tatacaggga gctactactt tgatggaacc 841 ctctacaaca aagggttcaa gggcgaaaag gaccactggc ggagccgacg caccatctgt 901 gtcaaaaccc acgagagtgg caggagggag attgagatgt ttgattcagc catcctgcta 961 atccqqaacc catacaqqtc cctggtggca gaattcaaca gaaaatgtgc cgggcacctg 1021 qqatatqcaq ctqaccqcaa ctqqaaqaqc aaagagtggc cggactttgt caacagctac 1081 gcctcgtggt ggtcctcgca cgtcctggac tggctcaagt acgggaagcg gctgctggtg 1141 gtgcactacg aggagctgcg gcgcagcctg gtgcccacgt tacgggagat ggtggccttc 1201 ctcaacqtgt ctgtgagcga ggagcggctg ctctgcgtgg agaacaacaa ggagggcagc 1261 ttccggcggc gcggccggcg ctcccacgac cctgagccct tcaccccgga gatgaaagac 1321 ttgatcaatg gctacatcg gacggtggac caagccctgc gtgaccacaa ctggacgggg 1381 ctgcccaggg agtatgtgcc cagatgatag gcctggccca cgccgccgcc cccgctgagt 1441 gacgcaatcg caccacgggg ctgcgctccc cactctgatg ctcaggcccg tggcctcact 1501 gggacgaacg gtgggtgggg ggctcaccct ggtgctgcct cccgcacaag gagacctgga 1561 cacaacagac acacatcaca aggcgaacac aaatggacac acatacctgg ccacgaaccc 1621 acacetecte agacacteag acaceaetee aggeteatag eccegtettg atgeagagaa 1681 gccacccacg tggggtgtgc caggcacccc cagatacaaa tgcagccacg cacagacgta 1741 acacacaggt gccaggccgt gtgctcctgg aggctggctg gctgtctctc tcacacagat 1801 acacqtqcqc tccctqqqat ccqqqaqqcc ctqqqcttcc tqtqtqtaqc cctqqcataq 1861 acttgctcgt cagggtgttc gactctggga tgctgggccg ggcagacatt tatgctctga 1921 gcagcaagga ccattgggat ggaggtgggc acaaagactg ctgcttccag ggtgtgcggc 1981 cctqqccqtq tqtctqacat cccataaatq tqtqtqtqtq qtqactacqq qcaccacaaa 2101 aggactgccc cccaccccgc ccaagccaat gcagacactg acctcaagac cagccacacc 2161 aagtetgtge actgeecace acceetteg acceetecta caccatgete etgggeetgg 2221 ggcctccttg cctggcttct tcctccagct atcctcaggg aagctggaga ttcaagtcat 2281 gccccagaag caaggtgtaa gcatgggggc gtggggaaga gggcaggagc tgaagttcca 2341 ggggagccca ggacccctgg ggacctgtac actagcagga tgggtcttgg gagttggcat 2401 ggggagattg taagggctgt atctccgcga ttactggtac tggtgctcca cccatagggg 2461 ccctgttctc tctggggtcc cacactgggg tcatgagagc tagttctcat gagagcagga 2521 cccctaggtc tggtcctgcc agcctgggat gggggtcaat gtatgctaca catcttgggc 2581 tcacaagtga qqctqtqqaq qccaqqagqt tcaaqtatqq agggtaggag tgcaacccaq 2641 gagcaacgct caaggtggcc gagatgcagg gaggattgtg gcgatccggt tgcacactgg 2701 cctctgtgga ggtcagggag gtgggaaggt gggtgggcct gaactcaggc agcccttcag 2761 tggcttcttc aggcctagga taaccctggc caggccccat ggtcctgctc tgggtcagct 2821 qtcagggaga cctcccttca qcctgagcca agctgcattg ggaggagaag gcagggtgag 2881 ggatctggga gagcaggatg ttgagatatc ctgaccctct gcttcctagc ctgatgatgg 2941 ggaacacaga gccacccgct atgagcacag gccgggggtg gaaggcccag tagagctgcc 3001 cttccccaga tggaagcttc ctccaacctt cagtttggcc agtgggtggg tttgggcatc 3061 tgtgcccctg gtggcctcca aatccttcct gggattcccc caaacccaca ctgatgcgaa 3121 gctgttcttg acceteccat agcateaact cagttcaget caaagetaga aacaccaagg 3181 tetttgaaaa tgtttattgg etcaggatag ettcagcate caatetgega aggeeeetga 3241 tgqaqctatt tcttcctaqt atcaggagca attgagaccc ctcaggagtt tccacactgc

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3301 tgtggacggg atagccaggg cagacctgca tcccagaqca agcctagatc cttqaaaqca
    3361 atgtcaggcg tggcagtggg gaagccatgg aggctgggaa gggcaaagtg tggagagagg
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    3481 ttggattggt acattcactg ctgccaggta cccagagatg tcatctctct ccaggtagag
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    3841 totoatotac agotottago ttotoagtaa agaggatact otoccaacca gocaaggaat
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    3961 gaacagagaa accetgactg tetacagatt ecaaetttaa tagcaageet gtggagttae
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    4441 atqtqatqac tccaaaqqqq ttcatttcac tataqqqaqa aattgattcg tctgacttcc
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    4681 tggtaggtct tgctgattgc atccacattc attcctccat ttcctttccc agcaaccccc
    4741 catttctqtt tqqqaattca tatgatctgc cagagcctct ccaggggtgg aatacatgtc
    4801 ctggactaag ccaatgacac cttccatctt tccagctatg gtgactgggt taggtatgat
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    5041 ggagagttga gaggcagaaa gagatcaagt ccttggtgaa attgtttaag tacaggatca
    5101 agentiagett gagattggen ttachtetgg antitttttg ttatatgage taataaatte
    5161 tqtttattqt tt
//
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S NCBI	NCBI %		S Nucleotide			
PubMed Nucle	otide Protein	Genome	Structure	PMC	Taxonomy	OMIM Boo
Search Nucleotide	for				Go Clear	
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Display default	▼ Show: 20	▼ Send to	File	- ▼		

☐ 1: AA426220. zv84d01.s1 Soares...[gi:2107753]

Links

IDENTIFIERS

dbEST Id: EST name:

1070767 zv84d01.s1

GenBank Acc: AA426220 GenBank gi:

2107753

CLONE INFO

Clone Id:

IMAGE: 760321 (3')

Source:

IMAGE Consortium, LLNL

DNA type:

CDNA

PRIMERS

Sequencing:

-41m13 fwd. ET from Amersham

PolyA Tail:

Unknown

SEQUENCE

AAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGTAAGGCCAATC TCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTTCTGCC TCTCAACTCTCCCTTCAGTGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGCC CAAGGTGGTCATCATAAAGACCCAGGAATACTACTACCTTTTTCACATTCAACAGGGGAA

TTAAAACAGCTTCTACCCAGCAT

Quality:

High quality sequence stops at base: 194

Entry Created: May 19 1997 Last Updated:

Oct 16 1997

COMMENTS

This clone is available royalty-free through LLNL; contact

the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name:

Soares_total_fetus Nb2HF8 9w

Organism:

Homo sapiens Develop. stage: 8-9 weeks

Lab host:

DH10B

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I

R. Site 2:

Eco RI

Description:

1st strand cDNA was prepared from mRNA obtained from pooled 8-9 week (total) fetus material with a Not I - oligo(dT)

primer [5'

Double-stranded cDNA was ligated to Eco RI adaptors

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization, and was constructed by

Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: WashU-Merck EST Project 1997

Authors: Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost

,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising

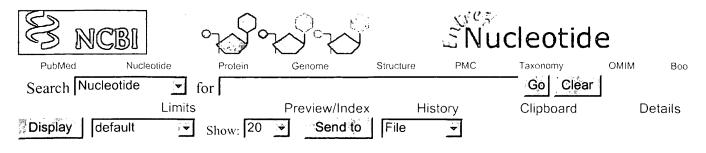
,B., White,Y., Wylie,T., Waterston,R., Wilson,R.

Year: 1997

Status: Unpublished

MAP DATA

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☐ 1: AI915364. wd38f08.x1 Soares...[gi:5635219]

Links

IDENTIFIERS

 dbEST Id:
 3037330

 EST name:
 wd38f08.x1

 GenBank Acc:
 AI915364

 GenBank gi:
 5635219

CLONE INFO

Clone Id: IMAGE:2330439 (3')

Source: NCI Insert length: 459 DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: Unknown

SEQUENCE

TTTTTTTTTAAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGT AAGGCCAATCTCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATC TCTTTCTGCCTCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGA TCCCAAGGCCCAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTCACATTC AACAGGGGAATTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGG

Quality: High quality sequence stops at base: 272

Entry Created: Jul 28 1999 Last Updated: Dec 17 1999

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares_NFL_T_GBC_S1

Organism: Homo sapiens

Organ: pooled Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: Equal amounts of plasmid DNA from three normalized libraries

(fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087,

682632-687239, 726408-728711, and 729096-731399. Subtraction

by Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

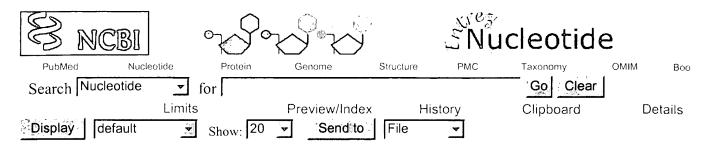
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

MAP DATA

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☐ 1: Z39512. HSC1BH062 normali...[gi:562704]

Links

IDENTIFIERS

dbEST Id: 64601
EST name: HSC1BH062
GenBank Acc: Z39512
GenBank gi: 562704

CLONE INFO

Clone Id: c-1bh06 (3')

DNA type: cDNA

PRIMERS

Sequencing: (-21)M13 universal

PolyA Tail: Unknown

SEQUENCE

AAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGTAAGGCCAATC
TCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTTCTGCC
TCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGCC
CAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTTCACATTCAACAGGGGAA
TTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGGCTTCATTGTC

 ${\tt TTAACTCACANGATCATACCTAACCCAGTCACCATAGCTGGAAAGATG}$

Entry Created: Oct 31 1994 Last Updated: Oct 31 1994

COMMENTS

Single read. 25 T removed at sequence 5'end

Genexpress_library_idt: C; Genexpress sequence idt:

a1c-1bh06

LIBRARY

Lib Name: normalized infant brain cDNA

Organism: Homo sapiens

Sex: Female Organ: brain

Tissue type: total brain
Develop. stage: 3 months old
Vector: lafmid BA
R. Site 1: HindIII
R. Site 2: NotI

Description: sex=Female; dev_stage=3 months old; isolate=muscular atrophy

patient; tissue_type=total brain; total mRNA was oligo-(dT)
primed and directionally cloned 5' -> 3' into the HindIII ->
NotI sites of the lafmid BA vector. Clone library from

B.Soares, Psychiatry Dept. Columbia University, USA. Normalization_method: Bento Soares, P.N.A.S in press

SUBMITTER

Name: Genethon

Lab: Genexpress-Genethon

Institution: Genethon Centre de recherche sur le Genome Humain

Address: 1, rue de l'Internationale, BP60 91002 EVRY Cedex, FRANCE

Tel: 33169472800 Fax: 33160778698

E-mail: genexpress@genethon.fr

CITATIONS

Medline UID: <u>95277534</u>

Title: IMAGE: molecular integration of the analysis of the human

genome and its expression

Authors: Auffray, C., Behar, G., Bois, F., Bouchier, C., da Silva, C.,

Devignes, M.D., Duprat, S., Houlgatte, R., Jumeau, M.N., Lamy, B.

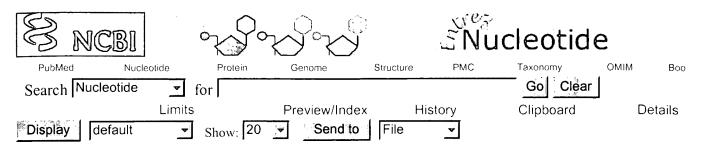
, Lorenzo, F., Mitchell, H., Mariage-Samson, R., Pietu, G.,

Pouliot, Y., Sebastiani-Kabaktchis, C., Tessier, A.

Citation: C. R. Acad. Sci. III, Sci. Vie 318 (2): 263-272 1995

MAP DATA

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☐ 1: N63086. yz32e07.s1 Soares...[gi:1210915]

Links

IDENTIFIERS

 dbEST Id:
 468837

 EST name:
 yz32e07.s1

 GenBank Acc:
 N63086

 GenBank gi:
 1210915

 GDB Id:
 3903300

CLONE INFO

Clone Id: IMAGE:284772 (3')

Source: IMAGE Consortium, LLNL

Insert length: 724 DNA type: cDNA

PRIMERS

Sequencing: m13 -40 forward

PolyA Tail: Unknown

SEQUENCE

AAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGTAAGGCCAATC TCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTCTGCC TCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGCC CAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTCACATTCAACAGGGGAA TTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGGCTTCATTGTC

TTAACTCACAAGATCATACCTAACCCCAGTCACCATAAGCTGGAAAGATGGA

Quality: High quality sequence stops at base: 285

Entry Created: Feb 23 1996 Last Updated: Jan 30 1997

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares_multiple_sclerosis_2NbHMSP

Organism: <u>Homo sapiens</u>

Sex: male

Tissue type: multiple sclerosis lesions

Develop. stage: Age 46

Lab host: DH10B (ampicillin resistant)

Vector: pT7T3D (Pharmacia) with a modified polylinker V TYPE:

phagemid

R. Site 1: Not I R. Site 2: Eco RI

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector

(Pharmacia). Library went through one round of normalization

to a Cot = 5. Library constructed by Bento Soares and

M.Fatima Bonaldo. RNA from 4 multiple sclerosis lesions from one patient was kindly provided by Dr. Kevin G. Becker

(NINDS/NIH).

SUBMITTER

Name: Wil

Wilson RK

Institution: Address:

Washington University School of Medicine

Tel:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108 314 286 1800

Tel: Fax:

314 286 1810

E-mail:

est@watson.wustl.edu

CITATIONS

Medline UID:

97044478

Title:

Generation and analysis of 280,000 human expressed sequence

tags

Authors:

Hillier, L., Lennon, G., Becker, M., Bonaldo, M.F., Chiapelli, B., Chissoe, S., Dietrich, N., DuBuque, T., Favello, A., Gish, W., Hawkins, M., Hultman, M., Kucaba, T., Lacy, M., Le, M., Le, N., Mardis, E., Moore, B., Morris, M., Parsons, J., Prange, C., Rifkin, L., Rohlfing, T., Schellenberg, K., Soares, M.B., Tan, F., Thierry-Meg, J., Trevaskis, E., Underwood, K., Wohldmann, P.,

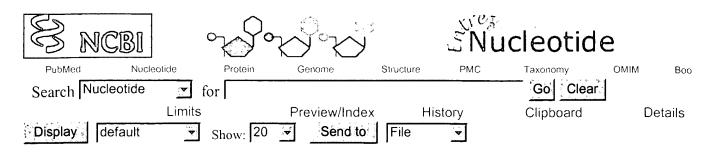
Waterston, R., Wilson, R., Marra, M.

Citation:

Genome Res. 6 (9): 807-828 1996

MAP DATA

<u>Disclaimer</u> | <u>Write to the Help Desk</u> <u>NCBI</u> | <u>NLM</u> | <u>NIH</u>



☐ 1: AI095108. qa14h11.x1 NCI_CG...[gi:3434084]

Links

IDENTIFIERS

 dbEST Id:
 1853495

 EST name:
 qa14h11.x1

 GenBank Acc:
 AI095108

 GenBank gi:
 3434084

CLONE INFO

Clone Id: IMAGE:1686789 (3')

Source: NCI Insert length: 724 DNA type: cDNA

PRIMERS

Sequencing: -40ml3 fwd. ET from Amersham

PolyA Tail: Unknown

SEQUENCE

TAAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGTAAGGCCAAT CTCAAGCAAGGCTTGATCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTTCTGC CTCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGC CCAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTCACATTCACATCACAGGGA ATTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGGCTTCATTGT CTTAACTCACAAGATCATACCTAACCCAGTCACCATAGCTGGAAAGATGGAAGGTGTCAT

TGGCTTAGTCCAGGACATGTATTCCACCCCTGGAGAG

Quality:

High quality sequence stops at base: 396

Entry Created: Aug 18 1998 Last Updated: Oct 1 1998

COMMENTS

Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld

M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima

Bonaldo, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Brn23
Organism: Homo sapiens

Organ: brain

Tissue type: glioblastoma (pooled)

Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

[5'

Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute / National Institute of

Neurological Disorders and Stroke, Brain Tumor Genome

Anatomy Project (CGAP/BTGAP), Tumor Gene Index

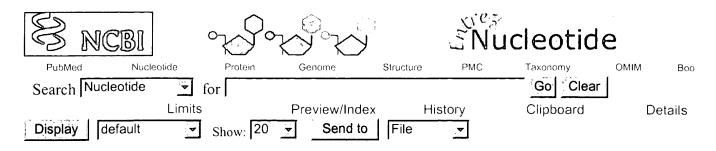
Authors: NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1998

Status: Unpublished

MAP DATA

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1: AI056703. oy53e04.x1 NCI_CG...[gi:3330569]

Links

IDENTIFIERS

dbEST Id:

1796154

EST name: GenBank Acc: oy53e04.x1 AI056703

GenBank gi:

3330569

CLONE INFO

Clone Id:

IMAGE:1669566 (3')

Source:

NCI

DNA type:

cDNA

PRIMERS

Sequencing:

-40ml3 fwd. ET from Amersham

PolyA Tail:

Unknown

SEQUENCE

TAAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGTAAGGCCAAT CTCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTTCTGC CTCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGC CCAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTCACATTCAACAGGGGA ATTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGGCTTCATTGT CTTAACTCACAAGATCATACCTAACCCAGTCACCATAGCTGGAAAGATGGAAGGTGTCAT TGGCTTAGTCCAGGACATGTATTCCACCCCTGGAGAGGCTCTTGGCAGATCATATGAATTC CCAAACAGAAATGGGGGGTT

Entry Created:

Jul 20 1998

Last Updated:

Jul 20 1998

COMMENTS

Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld

M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima

Bonaldo, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name:

NCI_CGAP_Brn23

Organism:

Homo sapiens

Organ:

brain

Tissue type:

glioblastoma (pooled)

Lab host:

DH10B

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I

R. Site 2:

Eco RI

Description:

1st strand cDNA was primed with a Not I - oligo(dT) primer

3']; double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library is normalized, and was constructed by Bento Soares and M.Fatima

Bonaldo.

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome

Anatomy Project (CGAP/BTGAP), Tumor Gene Index

Authors:

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

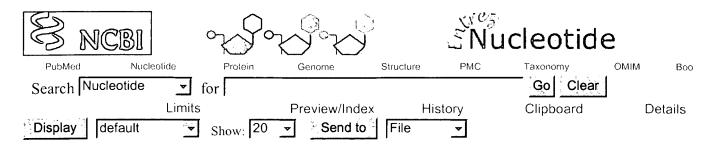
1998

Status:

Unpublished

MAP DATA

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☐ 1: BE326640. hr62f06.x1 NCI_CG...[gi:9200416]

Links

IDENTIFIERS

 dbEST Id:
 5303724

 EST name:
 hr62f06.x1

 GenBank Acc:
 BE326640

 GenBank gi:
 9200416

CLONE INFO

Clone Id: IMAGE:3133091 (3')

Source: NCI DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: Unknown

SEQUENCE

Entry Created: Jul 14 2000 Last Updated: Jul 14 2000

COMMENTS

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,

Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL, send

email to: info@image.llnl.gov

LIBRARY

Lib Name: NCI_CGAP_Kid11
Organism: Homo sapiens

Organ: kidney Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI Description: Plasmid DNA from the normalized library NCI_CGAP_Kid3 was

prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library

(cloneIDs 1322376-1323911, 1456007-1456775, and

1500552-1502855). Subtraction by Bento Soares and M. Fatima

Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

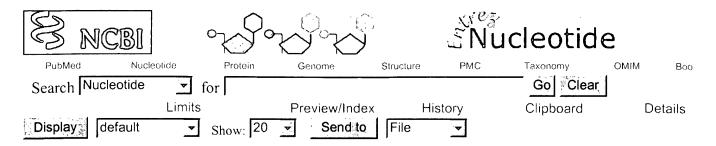
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

MAP DATA

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☐ 1: Al359305. qy29c09.x1 NCI_CG...[gi:4110926]

Links

IDENTIFIERS

dbEST Id:

2143123

EST name: GenBank Acc: qy29c09.x1

GenBank gi:

AI359305 4110926

CLONE INFO

Clone Id:

IMAGE:2013424 (3')

Source:

NCI

Insert length:

730

DNA type:

CDNA

PRIMERS

Sequencing:

-40UP from Gibco

PolyA Tail:

Unknown

SEQUENCE

AAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAGTCCAGAGGTAAGGCCAATC TCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTTCTGCC TCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGCC CAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTCACATTCAACAGGGGAA TTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGGCTTCATTGTC TTAACTCACAAGATCATACCTAACCCAGTCACCATAGCTGGAAAGATGGAAGGTGTCATT GGCTTAGTCCAGGACATGTATTCCACCCCTGGAGAGGCTCTGGCAGATCATATGAATTCC

Entry Created:

Jan 6 1999

Last Updated:

Feb 15 1999

COMMENTS

Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld

M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima

Bonaldo, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name:

NCI_CGAP_Brn23

Organism:

Homo sapiens

Organ:

brain

Tissue type: glioblastoma (pooled)

Lab host:

DH10B

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I

R. Site 2:

Eco RI

Description:

1st strand cDNA was primed with a Not I - oligo(dT) primer

[5]

Bonaldo.

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome

Anatomy Project (CGAP/BTGAP), Tumor Gene Index

Authors:

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

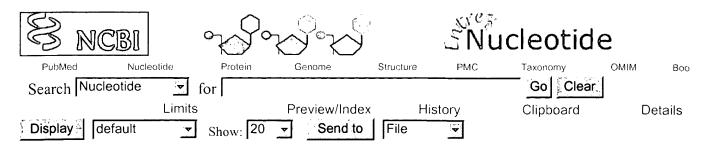
1998

Status:

Unpublished

MAP DATA

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☐ 1: AI421276. tf19g05.x1 NCl CG...[gi:4267207]

Links

IDENTIFIERS

 dbEST Id:
 2229032

 EST name:
 tf19g05.x1

 GenBank Acc:
 AI421276

 GenBank gi:
 4267207

CLONE INFO

Clone Id: IMAGE:2096696 (3')

Source: NCI Insert length: 745 DNA type: CDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: Unknown

SEQUENCE

С

Quality: High quality sequence stops at base: 451

Entry Created: Feb 10 1999 Last Updated: Mar 28 1999

COMMENTS

Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld

M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima

Bonaldo, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Brn23
Organism: Homo sapiens

Organ: brain

Tissue type: glioblastoma (pooled)

Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

[5'

Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute / National Institute of

Neurological Disorders and Stroke, Brain Tumor Genome

Anatomy Project (CGAP/BTGAP), Tumor Gene Index

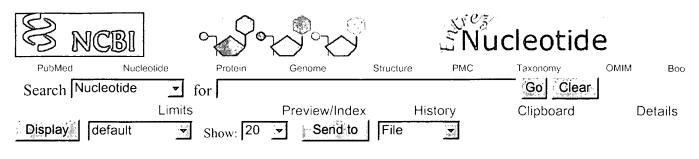
Authors: NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1998

Status: Unpublished

MAP DATA

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☐ 1: AW008726. ws72c12.x1 NCI CG...[gi:5857504]

Links

IDENTIFIERS

dbEST Id:

3137714

EST name:

ws72c12.x1 AW008726

GenBank Acc:

GenBank gi:

5857504

CLONE INFO

Clone Id:

IMAGE: 2502742 (3')

Source:

NCI 679

Insert length: DNA type:

CDNA

PRIMERS

Sequencing:

-40UP from Gibco

PolyA Tail:

Unknown

SEQUENCE

AAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGTAAGGCCAATC TCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTTCTGCC TCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGCC CAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTTCACATTCAACAGGGGAA TTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGGCTTCATTGTC TTAACTCACAAGATCATACCTAACCCAGTCACCATAGCTGGAAAGATGGAAGGTGTCATT GGCTTAGTCCAGGACATGTATTCCACCCCTGGAGAGGCTCTGGCAGATCATATGAATTCC

CAAGACCTACC

Quality:

High quality sequence stops at base: 465

Entry Created:

Last Updated:

Sep 9 1999

Mar 8 2000

COMMENTS

Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld

M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima

Bonaldo, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name:

NCI_CGAP_Brn23 Homo sapiens

Organism: Organ:

brain

Tissue type:

glioblastoma (pooled)

Lab host:

DH10B

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I Eco RI

R. Site 2: Description:

1st strand cDNA was primed with a Not I - oligo(dT) primer

[5]

Bonaldo.

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome

Anatomy Project (CGAP/BTGAP), Tumor Gene Index

Authors:

NCI/NINDS-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

1998

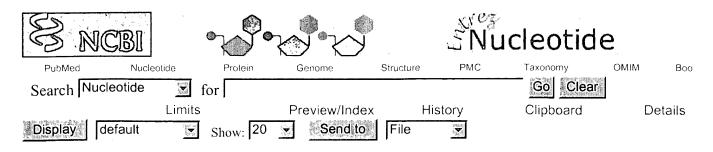
Status:

Unpublished

MAP DATA

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1: Al685123. wc67g02.x1 NCI_CG...[gi:4896417]

Links

IDENTIFIERS

 dbEST Id:
 2584900

 EST name:
 wc67g02.x1

 GenBank Acc:
 AI685123

 GenBank gi:
 4896417

CLONE INFO

Clone Id: IMAGE:2323730 (3')

Source: NCI Insert length: 766 DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: Unknown

SEQUENCE

TAAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTCCAGAGGTAAGGCCAAT CTCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGACTTGATCTCTTTCTGC CTCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTGGTCATGATCCCAAGGC CCAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTTCACATTCAACAGGGGA ATTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTCCCTCTGGCTTCATTGT CTTAACTCACAAGATCATACCTAACCCAGTCACCATAGCTGGAAAGATGAAGGTGTCAT TGGCTTAGTCCAGGACATGTATTCCACCCCTGGAGAGGCTCTGGCAGATCATATGAATTC CCAAACAGAAATGGGGGGTTGCTGGGAAAGGAAGATGAATGCACAATCA

GCAAGACCTACCACTTC

Quality: High quality sequence stops at base: 412

Entry Created: May 27 1999 Last Updated: Dec 17 1999

COMMENTS

Life Technologies catalog #: 11548-013

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Pan1
Organism: Homo sapiens
Organ: pancreas

Tissue type: adenocarcinoma

Lab host: DH10B

Vector: pCMV-SPORT6

R. Site 1: SalI

R. Site 2: NotI

Description: Cloned unidirectionally. Primer: Oligo dT. Average insert

size 1.72 kb. Life Technologies catalog #: 11548-013

SUBMITTER

¥

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

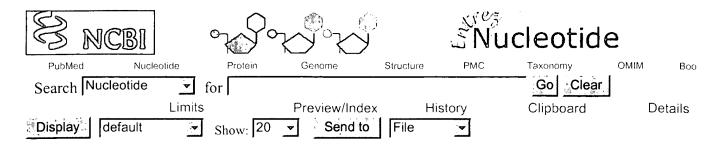
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

MAP DATA

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☐ 1: AW294233. UI-H-BI2-ahc-f-12...[gi:6700869]

Links

IDENTIFIERS

dbEST Id:

3677341

EST name:

UI-H-BI2-ahc-f-12-0-UI.s1

GenBank Acc:

AW294233

GenBank gi:

6700869

CLONE INFO

Clone Id:

IMAGE:2726542 (3')

Source:

NCI

DNA type:

CDNA

PRIMERS

Sequencing:

M13 Forward

PolyA Tail:

yes

SEQUENCE

TTTTTTTTTTTTTAAACAATAAACAGAATTTATTAGCTCATATAACAAAAAAAGTC CAGAGGTAAGGCCAATCTCAAGCAAGGCTTGATCCTGTACTTAAACAATTTCACCAAGGA CTTGATCTCTTTCTGCCTCTCAACTCTCCCTTCAGTGGTGTCAGCTTCACGTGATTCCTG GTCATGATCCCAAGGCCCAAGGTGGTCATCATAAAGACCCAGGAATACTACCTTTTT CACATTCAACAGGGGAATTAAAACAGCTTCTACCCAGCATTCCCATAAGTCTTGAAATTC CCTCTGGCTTCATTGTCTTAACTCACAAGATCATAACCCAGTCACCATAGCTGGAA AGATGGAAGGTGTCATTGGCTTAGTCCAGGACATGTATTCCACCCCTGGAGAGGCTCTGG CAGATCATATGAATTCCCAAACAGAAATGGGGGGTTGCTGGGAAAGGAAATGGAGGAATG

Entry Created:

Jan 14 2000

Last Updated:

Jan 16 2000

COMMENTS

The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of

first strand cDNA and therefore this may represent a

bonafide poly A tail. cDNA Library Preparation: M.B. Soares

Lab Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E.

Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html The following repetitive elements were found in this cDNA

sequence: 1-27, >AT_rich#Low complexity

AATGTGGATGCAATCAGCAAGAACTACCACTT

LIBRARY

Lib Name:

NCI CGAP Sub4

Organism:

Homo sapiens

Tag Seq: Tag Tissue: ATATC brain

Tag Lib:

NCI CGAP Brn23

Lab host:

DH10B (Life Technologies)

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: The NCI_CGAP_Sub4 library is a subtracted library derived

from the NCI_CGAP_Sub2 library which is a subtracted library

derived from the NCI_CGAP_Sub1 library, which is a

subtracted library derived from BI. BI constitutes a mixture

of 21 normalized or subtracted NCI_CGAP libraries:

NCI_CGAP_Co4, NCI_CGAP_Pr22, NCI_CGAP_Pr28, NCI_CGAP_Co10, NCI_CGAP_Co16, NCI_CGAP_Kid5, NCI_CGAP_Kid12, NCI_CGAP_Kid3, NCI_CGAP_Kid11, NCI_CGAP_Lym2, NCI_CGAP_Br2, NCI_CGAP_Co8, NCI_CGAP_CLL1, NCI_CGAP_Lei2, NCI_CGAP_Brn23, NCI_CGAP_Lu5, NCI_CGAP_Lu24, NCI_CGAP_Lu19, NCI_CGAP_GC4, NCI_CGAP_GC6,

NCI_CGAP_Brn25. These 21 libraries were pooled and a

single-stranded DNA preparation of the resulting mixture was used as a tracer in a subtractive hybridization with a driver whose composition is detailed below: NCI_CGAP_Kid3 pool 1 : LLAM 3334-3337, 3682-3683, 3798-3803 (IMAGE

CloneIDs 1322376-1323911, 1456008-1456775, 1500552-1502855) NCI CGAP Kid5 pool 1 : LLAM 3338-3342, 3722-3725, 3776-3778

(IMAGE CloneIDs 1323912-1325831, 1471368-1472903, 1492104-1493255) NCI_CGAP_Lu5 pool 1 : LLAM 3575-3582, 3851-3854 (IMAGE CloneIDs 1414920-1417991, 1520904-1522439) NCI_CGAP_GC4 pool 1 : LLAM 3164-3167, 3716-3720, 3733-3735

(IMAGE CloneIDs 1257096-1258631, 1469064-1470983, 1475592-1476743) NCI_CGAP_Pr22 pool 1 : LLAM 2457-2459, 2758-2759, 3062-3068 (IMAGE CloneIDs 985608-986759, 1101192-1101959, 1217928-1220615) NCI_CGAP_Col0 pool 1 : LLAM 2644-2653, 2871-2872 (IMAGE CloneIDs 1057416-1061255, 1144584-1145351) Subtraction was performed as previously

described [Bonaldo, Lennon & Soares (1996): Normalization and Subtraction: Two Approaches To Facilitate Gene

Discovery. Genome Research 6, 791-806.]

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: <u>cgapbs-r@mail.nih.gov</u>

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

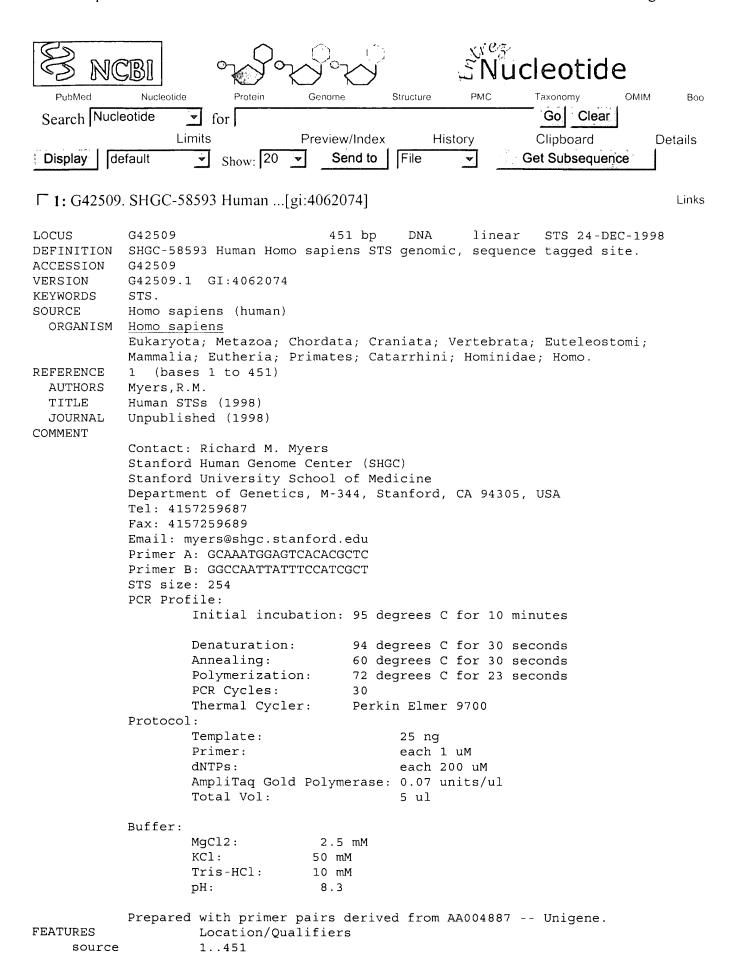
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

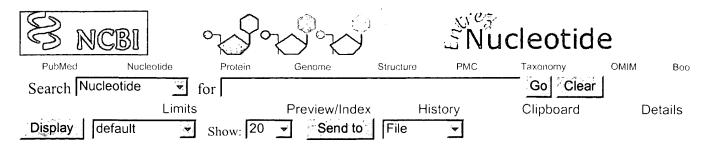
MAP DATA

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/organism="Homo sapiens"
                    /mol type="genomic DNA"
                    /db_xref="taxon:9606"
                    /map="11"
                    /clone lib="Human"
    STS
                    49..302
    primer bind
                    49..68
    primer bind complement(283..302)
BASE COUNT
               134 a 77 c 97 g
                                       142 t 1 others
ORIGIN
       1 gcagaggcct ccacttttta tttcagttgt actcatctgt cccactgtgc aaatggagtc
      61 acacgeteae teaattetga gaggeetgge aagnaaagag aaaagatgee cagageagte
     121 tgttagagtt gcattctcag actaatatct ttacagtctt gagaaatcac tgtcagggtt
     181 tatttaaaat gcagattttt gaaggataaa ttttacgact aattttttt aataaactat
     241 gcaggattgt tatttagaag atttgccaaa tttagagtct tcagcgatgg aaataattgg
     301 ccttcttgtc acagtcttct gtttataagt gggtaaagaa agttttcttt ccagaaaaat
     361 acagcagaaa atccgatggt tctgatagga gttaattgtg gagatgtgcc agagacagca
     421 gcttcgtgga tggtgacacc acaatgtctg t
//
```

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☐ 1: AA004887. zh90g01.s1 Soares...[gi:1447704]

Links

IDENTIFIERS

dbEST Id:

615134

EST name:

zh90g01.s1

GenBank Acc:

AA004887

GenBank gi:

1447704

GDB Id:

1328361

CLONE INFO

Clone Id:

IMAGE:428592 (3')

Source:

IMAGE Consortium, LLNL

DNA type:

CDNA

PRIMERS

Sequencing:

mob.REGA+ET

PolyA Tail:

Unknown

SEQUENCE

GCAGAGGCCTCCACTTTTTATTTCAGTTGTACTCATCTGTCCCACTGTGCAAATGGAGTC ACACGCTCACTCAATTCTGAGAGGCCTGGCAAGNAAAGAGAAAAGATGCCCAGAGCAGTC TGTTAGAGTTGCATTCTCAGACTAATATCTTTACAGTCTTGAGAAATCACTGTCAGGGTT GCAGGATTGTTATTTAGAAGATTTGCCAAATTTAGAGTCTTCAGCGATGGAAATAATTGG ACAGCAGAAAATCCGATGGTTCTGATAGGAGTTAATTGTGGAGATGTGCCAGAGACAGCA

GCTTCGTGGATGGTGACACCACAATGTCTGT

Quality:

High quality sequence stops at base: 429

Entry Created:

Jul 23 1996

Last Updated:

Jul 23 1996

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name:

Soares_fetal_liver_spleen_1NFLS_S1

Organism:

Homo sapiens

Sex:

male

Organ:

Liver and Spleen

Lab host:

Develop. stage: 20 week-post conception fetus

Vector:

DH10B (ampicillin resistant)

R. Site 1:

pT7T3D (Pharmacia) with a modified polylinker

R. Site 2:

Pac I

Eco RI

Description:

This is a subtracted version of the original Soares fetal

liver spleen 1NFLS library. 1st strand cDNA was primed with

a Pac I - oligo(dT) primer [5'

(Pharmacia), digested with Pac I and cloned into the Pac I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: Fax: 314 286 1800 314 286 1810

E-mail:

est@watson.wustl.edu

CITATIONS

Title:

The WashU-Merck EST Project

Authors:

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M., Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,

Wilson, R.: 1995

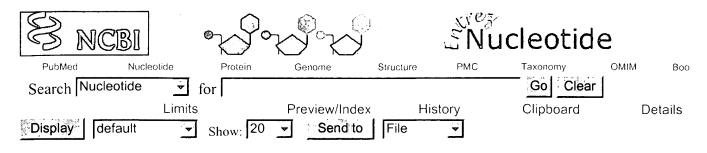
Year:

Status:

Unpublished

MAP DATA

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1: AA281006. zs94a06.s1 NCI_CG...[gi:1923895]

Links

IDENTIFIERS

 dbEST Id:
 922124

 EST name:
 zs94a06.s1

 GenBank Acc:
 AA281006

 GenBank gi:
 1923895

 GDB Id:
 5854830

CLONE INFO

Clone Id: IMAGE:705106 (3')

Source: IMAGE Consortium, LLNL

Insert length: 845 DNA type: cDNA

PRIMERS

Sequencing: -41m13 fwd. ET from Amersham

PolyA Tail: Unknown

SEQUENCE

TTTTTTTTTAATTAGCTGTTCTTGTCATATAGTTTTATTCCTTTATCTTTTTTTGAAC
ATTTTATACACCCTTATTTCAATGTTCCTTTTAGATCACTCTATTCTCTTGG
GCTTTGAATCTCCTTGTTTCTTGTATCTGCTGCCTCTCTTTGGGATACCTGGGAGTTTTT
CCTCTGACCTCGTCTTCAGTAGGAAATGATTTTCCATGAGAATCCTGGTTCCCCTGGATG

AGGACGGTGTCTCCTGGGGAGAATGTCCTGTT

Quality: High quality sequence stops at base: 252

Entry Created: Mar 17 1997 Last Updated: Aug 14 1997

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: NCI_CGAP_GCB1
Organism: Homo sapiens

Tissue type: germinal center B cell

Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: 1st strand cDNA was prepared from human tonsillar cells

enriched for germinal center B cells by flow sorting (CD20+, IgD-), provided by Dr. Louis M. Staudt (NCI), Dr. David Allman (NCI) and Dr. Gerald Marti (CBER). cDNA synthesis was

primed with a Not I - oligo(dT) primer

Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization, and was constructed by

Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

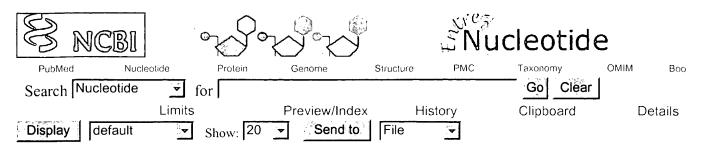
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

MAP DATA

<u>Disclaimer | Write to the Help Desk</u> <u>NCBI | NLM | NIH</u>



☐ 1: AA490819. aa49f05.s1 NCI CG...[gi:2219992]

Links

IDENTIFIERS

dbEST Id:

1135060

EST name:

aa49f05.s1

GenBank Acc: GenBank gi:

AA490819 2219992

CLONE INFO

Clone Id:

IMAGE:824289 (3')

Source:

NCI

DNA type:

CDNA

PRIMERS

Sequencing:

-41m13 fwd. ET from Amersham

PolyA Tail:

Unknown

SEQUENCE

TTTTTTTTCACTGTCACCATGAATTTAAATTTATTGAGTGCCCCACAAATGCTAGTCTA TTCTCAGTACATTTGATGAACACCATTTCTTTATCTCTAAAGGATGAGAGAATATTTGCT ACTATATTTTTTTTTGCTCATCACCCAGCCAGAATACAAATGGAACTCCTATGAATATT CTAAAGCATAATGAGGAAGGGGCTCCAGGCTAAATGCAAGTATCCTTGATTAATGTTTTC CCCACCACTGGGAATCACCCTCCCCGGTCCCCTGAAGCTTCCCCACAAGGTGCGGGGG

AAGCAGGAGAAAAAAAAGG

Entry Created:

Jun 25 1997

Last Updated:

Aug 15 1997

COMMENTS

Tissue Procurement: Louis M. Staudt, M.D., Ph.D., David

Allman, Ph.D., Gerald Marti, M.D.

cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima

Bonaldo, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name:

NCI CGAP GCB1

Organism:

Homo sapiens

Tissue type:

germinal center B cell

Lab host:

DH10B

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I

R. Site 2: Description: Eco RI 1st strand cDNA was prepared from human tonsillar cells enriched for germinal center B cells by flow sorting (CD20+, IgD-), provided by Dr. Louis M. Staudt (NCI), Dr. David Allman (NCI) and Dr. Gerald Marti (CBER). cDNA synthesis was primed with a Not I - oligo(dT) primer

Double-stranded cDNA was ligated to Eco RI adaptors

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization, and was constructed by

Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

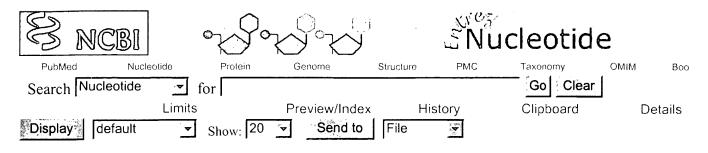
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

MAP DATA

<u>Disclaimer | Write to the Help Desk</u> <u>NCBI | NLM | NIH</u>



☐ 1: AA490870. aa48b07.s1 NCI_CG...[gi:2220043]

Links

IDENTIFIERS

 dbEST Id:
 1135111

 EST name:
 aa48b07.s1

 GenBank Acc:
 AA490870

 GenBank gi:
 2220043

CLONE INFO

Clone Id: IMAGE:824149 (3')

Source: NCI DNA type: cDNA

PRIMERS

Sequencing: -41m13 fwd. ET from Amersham

PolyA Tail: Unknown

SEQUENCE

TTTTTTTTCACTGTCACCATGAATTTAAATTTATTGAGTGCCCCACAAATGCTAGTCTA
TTCTCAGTACATTTGATGAACACCATTTCTTTATCTCTAAAGGATGAGAGAATATTTGCT
ACTATATATTTTTTTTGCTCATCACCCAGCCAGAATACAAATGGAACTCCTATGAATATT
CTAAAGCATAATGAGGAAGGGGCTCCAGGCTAAATGCAAGTATCCTTGATTAATGTTTTC
CCCACCACTGGGAATCACCCTCCCCCGCTCCCCTGAAGCTTCCCCACAAGGTGCGGGGG
AAGCAGGAAAAAAAAAGGAAGAGCCAAGGTATACTCCTGATAACGACACAAAAATTGAG

CAGTAGTAC

Quality: High quality sequence stops at base: 327

Entry Created: Jun 25 1997 Last Updated: Aug 15 1997

COMMENTS

Tissue Procurement: Louis M. Staudt, M.D., Ph.D., David

Allman, Ph.D., Gerald Marti, M.D.

cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima

Bonaldo, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_GCB1
Organism: Homo sapiens

Tissue type: germinal center B cell

Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I

R. Site 2:

Eco RI

Description:

1st strand cDNA was prepared from human tonsillar cells enriched for germinal center B cells by flow sorting (CD20+, IgD-), provided by Dr. Louis M. Staudt (NCI), Dr. David Allman (NCI) and Dr. Gerald Marti (CBER). cDNA synthesis was

primed with a Not I - oligo(dT) primer

Double-stranded cDNA was ligated to Eco RI adaptors

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization, and was constructed by

Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

Authors:

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

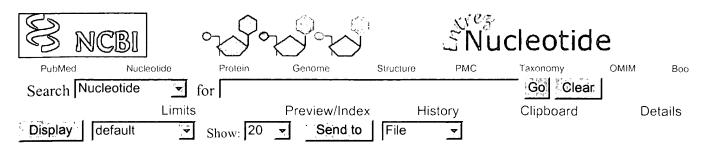
1997

Status:

Unpublished

MAP DATA

Disclaimer | Write to the Help Desk NCBI | NLM | NIH



☐ 1: N71063. za86a11.s1 Soares...[gi:1227643]

Links

IDENTIFIERS

dbEST Id: EST name: **477157** za86a11.s1

GenBank Acc: GenBank gi: N71063

GDB Id:

1227643 1244344

CLONE INFO

Clone Id:

IMAGE:299420 (3')

Source:

IMAGE Consortium, LLNL

DNA type: cDNA

PRIMERS

Sequencing:

m13 -40 forward

PolyA Tail:

Unknown

SEQUENCE

GATACTGCTCCCAGGTGTAAG

Quality:

High quality sequence stops at base: 344

Entry Created: Last Updated: Mar 14 1996 Mar 14 1996

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares_fetal_lung_NbHL19W

Organism: Homo sapiens

Organ: lung

Develop. stage: 19 weeks

Lab host: DH10B (ampicillin resistant)

Vector: pT7T3D (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector

(Pharmacia). Library went through one round of normalization

to a Cot = 5. Library constructed by Bento Soares and

 ${\tt M.Fatima}$ Bonaldo. This library was constructed from the same

fetus as the fetal heart library, Soares fetal heart

NbHH19W.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

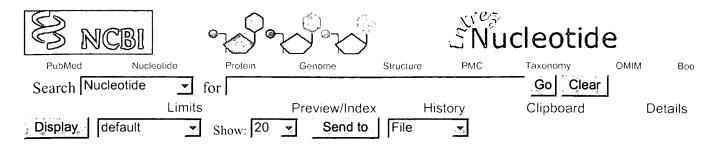
Wilson, R.

Year: 1995

Status: Unpublished

MAP DATA

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☐ 1: AA243738. zr68c12.s1 Soares...[gi:1874531]

Links

IDENTIFIERS

 dbEST Id:
 883127

 EST name:
 zr68c12.s1

 GenBank Acc:
 AA243738

 GenBank gi:
 1874531

 GDB Id:
 5562533

CLONE INFO

Clone Id: IMAGE:668566 (3')
Source: IMAGE Consortium, LLNL

DNA type: cDNA

PRIMERS

Sequencing: -41m13 fwd. ET from Amersham

PolyA Tail: Unknown

SEQUENCE

Quality: High quality sequence stops at base: 349

TCCCCATTCCAGTGGGCTGCTTTCTGGT

Entry Created: Mar 7 1997 Last Updated: Mar 7 1997

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (<u>info@image.llnl.gov</u>) for further

information.

LIBRARY

Lib Name: Soares_NhHMPu_S1
Organism: Homo sapiens
Organ: mixed (see below)

Tissue type: Pooled human melanocyte, fetal heart, and pregnant uterus

Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: Equal amounts of plasmid DNA from three normalized libraries

(melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and ss circles were made in vitro.

Following HAP purification, this DNA was used as tracer in a

subtractive hybridization reaction. The driver was

PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

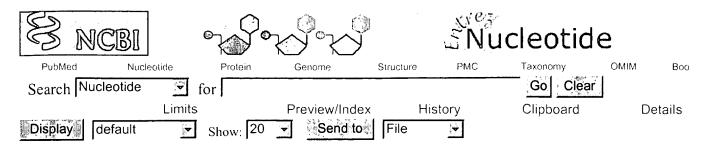
Wilson, R.

Year: 1995

Status: Unpublished

MAP DATA

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1: AA411711. zv16d08.s1 Soares...[gi:2069500]

Links

IDENTIFIERS

dbEST Id:

1054955

EST name:

zv16d08.s1 AA411711

GenBank Acc: GenBank gi:

2069500

GDB Id:

5976674

CLONE INFO

Clone Id:

IMAGE: 753807 (3')

Source:

IMAGE Consortium, LLNL

DNA type:

CDNA

PRIMERS

Sequencing:

-41m13 fwd. ET from Amersham

PolyA Tail:

Unknown

SEQUENCE

GCAGCCGCCTCCTAAGAACCTGCTGCTGGTCCCGGCAAGCCCAAGGAGCCAGCTGTGGT GCGCCGAGGAGCCTGTGGCAGCGCACATGCCAGAGGTAAAAAAACGACGCGG CGGAACAGAAGCTGGCATCTCCCCAGCCATCCTATGCAGCAGACGCCAACGACAGCAAGG CCGAGTACTCAGACGTCCTGGCCAAGCTGGCTTCCTGAACCGCCAGAGCCAGTGCGCTGG

ACGGTGCTCACCGCCCCGCTGCTGGACACCCAGTGAGCCGGAG

Entry Created:

May 2 1997

Last Updated:

May 17 1997

COMMENTS

This clone is available royalty-free through LLNL; contact

the IMAGE Consortium (info@image.llnl.gov) for further

information.

Possible reversed clone: polyT not found

LIBRARY

Lib Name:

Soares NhHMPu S1

Organism:

Homo sapiens

Organ:

mixed (see below)

Tissue type:

Pooled human melanocyte, fetal heart, and pregnant uterus

Lab host:

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: R. Site 2:

Not I

Description:

Eco RI

Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart

NbHH19W) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a

subtractive hybridization reaction. The driver was

PCR-amplified cDNAs from pools of 5,000 clones made from the

same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: Fax: 314 286 1800 314 286 1810

E-mail:

est@watson.wustl.edu

CITATIONS

Title:

WashU-Merck EST Project 1997

Authors:

Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R., Wilson, R.

Year:

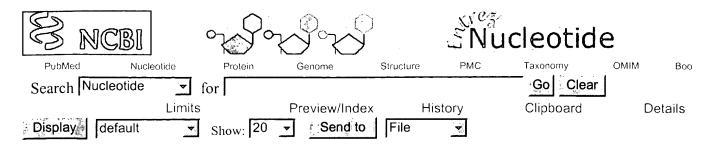
1997

Status:

Unpublished

MAP DATA

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☐ 1: N73808. yz80g07.s1 Soares...[gi:1231093]

Links

IDENTIFIERS

 dbEST Id:
 479945

 EST name:
 yz80g07.s1

 GenBank Acc:
 N73808

 GenBank gi:
 1231093

 GDB Id:
 3905244

CLONE INFO

Clone Id: IMAGE:289404 (3')

Source: IMAGE Consortium, LLNL

DNA type: cDNA

PRIMERS

Sequencing: m13 -40 forward

PolyA Tail: Unknown

SEQUENCE

 ${\tt AAAAAATGATTCAGTTAAAACAAAACAAAAGTTTAGATATTTTAG}$

Quality: High quality sequence stops at base: 144

Entry Created: Mar 19 1996 Last Updated: Mar 19 1996

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares multiple sclerosis_2NbHMSP

Organism: <u>Homo sapiens</u>

Sex: ${\sf male}$

Tissue type: multiple sclerosis lesions

Develop. stage: Age 46

Lab host: DH10B (ampicillin resistant)

Vector: pT7T3D (Pharmacia) with a modified polylinker V TYPE:

phagemid

R. Site 1: Not I R. Site 2: Eco RI

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

(Pharmacia). Library went through one round of normalization

to a Cot = 5. Library constructed by Bento Soares and M.Fatima Bonaldo. RNA from 4 multiple sclerosis lesions from one patient was kindly provided by Dr. Kevin G. Becker (NINDS/NIH).

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

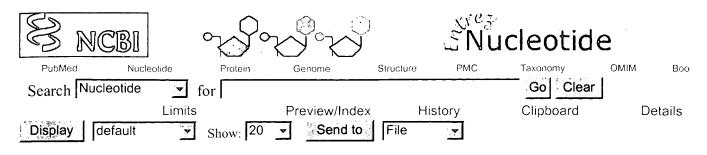
Wilson, R.

Year: 1995

Status: Unpublished

MAP DATA

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1: AI699181. tx61e11.x1 NCI_CG...[gi:4987081]

Links

IDENTIFIERS

 dbEST Id:
 2603719

 EST name:
 tx61e11.x1

 GenBank Acc:
 AI699181

 GenBank gi:
 4987081

CLONE INFO

Clone Id: IMAGE:2274092 (3')

Source: NCI Insert length: 2085 DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: Unknown

SEQUENCE

TGCAGGACCATAATCACATTTAAGCCACTGACT

Quality: High quality sequence stops at base: 200

Entry Created: Jun 3 1999 Last Updated: Dec 16 1999

COMMENTS

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,

Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: Life Technologies, Inc.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Ut1
Organism: Homo sapiens

Organ: uterus

Tissue type: well-differentiated endometrial adenocarcinoma, 7 pooled

tumors

Lab host: DH10B

Vector: pCMV-SPORT6

R. Site 1: SalI R. Site 2: NotI Description:

Cloned unidirectionally. Primer: Oligo dT. Average insert

size 1.75 kb. Life Technologies catalog #: 11538-014

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

Authors:

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

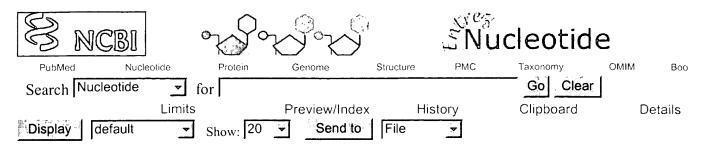
1997

Status:

Unpublished

MAP DATA

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☐ 1: R50866. yg62h08.s1 Soares...[gi:812768]

Links

IDENTIFIERS

 dbEST Id:
 220931

 EST name:
 yg62h08.s1

 GenBank Acc:
 R50866

 GenBank gi:
 812768

 GDB Id:
 410107

CLONE INFO

Clone Id: IMAGE: 37566 (3')

Insert length: 1925 DNA type: cDNA

PRIMERS

Sequencing: Promega -21m13

PolyA Tail: Unknown

SEQUENCE

ANTTCACACTGCATCTNGTTACTATAGTATTTCTAGGGCATA

Quality: High quality sequence stops at base: 251

Entry Created: May 18 1995 Last Updated: May 18 1995

COMMENTS

Insert Size: 1925

High quality sequence stops: 251 Source: IMAGE Consortium, LLNL This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for

further information.

LIBRARY

Lib Name: Soares infant brain 1NIB

Organism: Homo sapiens
Sex: female
Organ: whole brain

Develop. stage: 73 days post natal

Lab host: DH10B (ampicillin resistant)

Vector: Lafmid BA
R. Site 1: Not I
R. Site 2: Hind III

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

(Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafmid BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

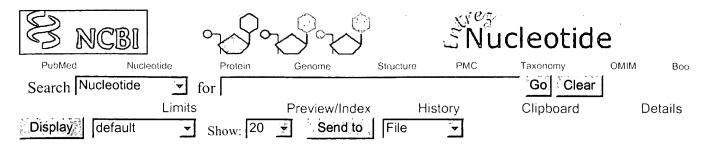
Wilson, R.

Year: 1995

Status: Unpublished

MAP DATA

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☐ 1: AI910763. wg96h11.x1 NCI CG...[gi:5630499]

Links

IDENTIFIERS

 dbEST Id:
 3032729

 EST name:
 wg96h11.x1

 GenBank Acc:
 AI910763

 GenBank gi:
 5630499

CLONE INFO

Clone Id: IMAGE:2379141 (3')

Source: NCI Insert length: 1427 DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: Unknown

SEQUENCE

TG

Quality: High quality sequence stops at base: 297

Entry Created: Jul 28 1999 Last Updated: Dec 20 1999

COMMENTS

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,

Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Kid11
Organism: Homo sapiens

Organ: kidney
Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I
R. Site 2: Eco RI

Description:

Plasmid DNA from the normalized library NCI CGAP Kid3 was prepared, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library

(cloneIDs 1322376-1323911, 1456007-1456775, and

1500552-1502855). Subtraction by Bento Soares and M. Fatima

Bonaldo.

SUBMITTER

Robert Strausberg, Ph.D. Name: E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

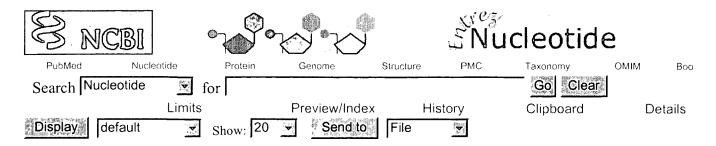
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap Authors:

Year: 1997

Unpublished Status:

MAP DATA

Disclaimer | Write to the Help Desk NCBI | NLM | NIH



1: AI351615. qr07f12.x1 Soares...[gi:4088821]

Links

IDENTIFIERS

 dbEST Id:
 2135429

 EST name:
 qr07f12.x1

 GenBank Acc:
 AI351615

 GenBank gi:
 4088821

CLONE INFO

Clone Id: IMAGE:1940207 (3')

Source: NCI DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: Unknown

SEQUENCE

Quality: High quality sequence stops at base: 310

Entry Created: Dec 30 1998 Last Updated: Dec 30 1998

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares_total_fetus Nb2HF8 9w

Organism: Homo sapiens
Develop. stage: 8-9 weeks

Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: 1st strand cDNA was prepared from mRNA obtained from pooled

8-9 week (total) fetus material with a Not I - oliqo(dT)

primer [5'

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went

through one round of normalization, and was constructed by Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

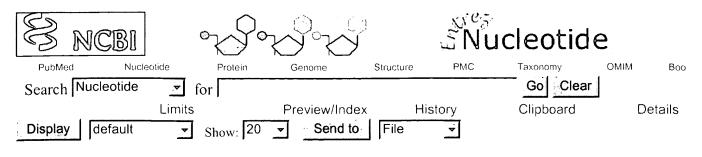
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

MAP DATA

<u>Disclaimer | Write to the Help Desk</u> <u>NCBI | NLM | NIH</u>



☐ 1: AA043474. zk54e09.s1 Soares...[gi:1521330]

Links

IDENTIFIERS

 dbEST Id:
 658848

 EST name:
 zk54e09.s1

 GenBank Acc:
 AA043474

 GenBank gi:
 1521330

 GDB Id:
 3760378

CLONE INFO

Clone Id: IMAGE:486664 (3')
Source: IMAGE Consortium, LLNL

DNA type: cDNA

PRIMERS

Sequencing: -40M13 fwd. from Amersham

PolyA Tail: Unknown

SEQUENCE

TGACAGCCATTGTACAAGTAA

Quality: High quality sequence stops at base: 301

Entry Created: Sep 4 1996 Last Updated: Sep 4 1996

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares_pregnant_uterus_NbHPU

Organism: Homo sapiens

Sex: female
Organ: uterus
Develop. stage: adult
Lab host: DH10B
Vector: pT7T3-Pac
R. Site 1: Not I
R. Site 2: Eco RI

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

 (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by M. Fatima Bonaldo.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

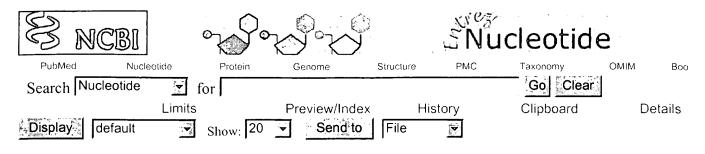
Wilson, R.

Year: 1995

Status: Unpublished

MAP DATA

<u>Disclaimer | Write to the Help Desk</u> <u>NCBI | NLM | NIH</u>



☐ 1: H08164. yl87e01.s1 Soares...[gi:872986]

Links

IDENTIFIERS

 dbEST Id:
 268044

 EST name:
 y187e01.s1

 GenBank Acc:
 H08164

 GenBank gi:
 872986

 GDB Id:
 417486

CLONE INFO

Clone Id: IMAGE: 44945 (3')

DNA type: cDNA

PRIMERS

Sequencing: Promega -21m13

PolyA Tail: Unknown

SEQUENCE

GGGATGTCACACAGTGACAGCCATTGTACAGGTAACGTTTAATGGACCTCCC

Quality: High quality sequence stops at base: 337

Entry Created: Jun 23 1995 Last Updated: Jun 23 1995

COMMENTS

High quality sequence stops: 337 Source: IMAGE Consortium, LLNL

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (<u>info@image.llnl.gov</u>) for further

information.

LIBRARY

Lib Name: Soares infant brain 1NIB

Organism: Homo sapiens
Sex: female

Organ: whole brain

Develop. stage: 73 days post natal

Lab host: DH10B (ampicillin resistant)

Vector: Lafmid BA
R. Site 1: Not I
R. Site 2: Hind III

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

double-stranded cDNA was ligated to Hind III adaptors (Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafmid BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

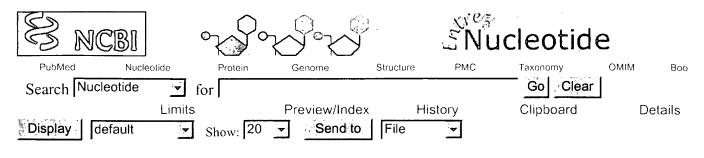
Wilson, R.

Year: 1995

Status: Unpublished

MAP DATA

<u>Disclaimer | Write to the Help Desk</u> <u>NCBI | NLM | NIH</u>



☐ 1: AI049699. an33d09.x1 Gessle...[gi:3298816]

Links

IDENTIFIERS

dbEST Id:

1789149

EST name: GenBank Acc: an33d09.x1 AI049699

GenBank gi:

3298816

CLONE INFO

Clone Id:

IMAGE:1700465 (3')

Source:

IMAGE Consortium, LLNL

DNA type:

CDNA

PRIMERS

Sequencing:

-40m13 fwd. ET from Amersham

PolyA Tail:

Unknown

SEQUENCE

TTTTTTTTTTTTTCTCTTTGAGTAACTTTATTTTGGAGGAGTTCCATAAGCATTAGGAAC TTGATTGATGCAATGTGATTGATTGATTGATTGATATACCAAAATTCATATGG

AAT

Entry Created:

Jul 9 1998

Last Updated:

Jul 9 1998

COMMENTS

This clone is available royalty-free through LLNL; contact

the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name:

Gessler Wilms tumor

Organism: Sex:

Homo sapiens

Lab host:

pooled (6) DH10B

Vector:

pSPORT1

R. Site 1:

SalI

R. Site 2:

NotI

Description:

RNA was prepared from a pool of 6 anonymous Wilms' tumor RNAs. RNA was prepared by acid-phenol, followed by one round of oligo dT selection. cDNA library preparation was with the BRL/Life Tech. Superscript Plasmid system. An oligo-dT NotI primer for first strand synthesis generated gcggccgccc(t)n at the 3' end of the clones. A 5' SalI adaptor was used with sequence 5'-gtcgacccacgcgtccg-3'. Resulting cDNAs were size selected (average size 2 kb), NotI digested, and ligated into NotI/SalI-cut pSPORT1. Library was constructed by Dr. Manfred Gessler.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: WashU-NCI human EST Project

Authors: Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost

,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R., Wilson

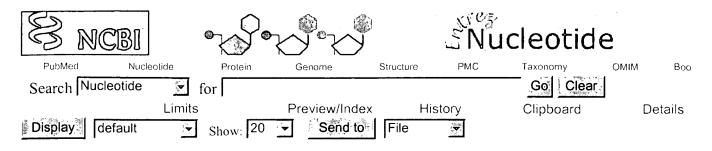
,R.

Year: 1997

Status: Unpublished

MAP DATA

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☐ 1: AW592865. hg04b07.x1 Soares...[gi:7280057]

Links

IDENTIFIERS

dbEST Id:

4025715

EST name: GenBank Acc: hq04b07.x1 AW592865

GenBank gi:

7280057

CLONE INFO

Clone Id:

IMAGE:2944597 (3')

Source:

NCI

DNA type:

CDNA

PRIMERS

Sequencing:

-40UP from Gibco

PolyA Tail:

Unknown

SEQUENCE

AATGACACCACTGTTGACAATGAAAAAAAAAACAGCATTTGATATTTTCCAGCTTTTT AAGTTAAAAATGATTCAGTTAAAACAAAACAAAAGTTTAGATATTTTAGCAAGAATCAC ATGCAATGTGATTGATTGATTGATTGATATATACCAAAATTCATATGGATGTCACA

TGG

Quality:

High quality sequence stops at base: 209

Entry Created:

Mar 22 2000

Last Updated:

Mar 22 2000

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name:

Soares_NFL_T_GBC_S1

Organism:

Homo sapiens

Organ:

pooled

Lab host:

DH10B

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I

R. Site 2:

Eco RI

Description:

Equal amounts of plasmid DNA from three normalized libraries

(fetal lung NbHL19W, testis NHT, and B-cell NCI CGAP GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs

from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

Authors:

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

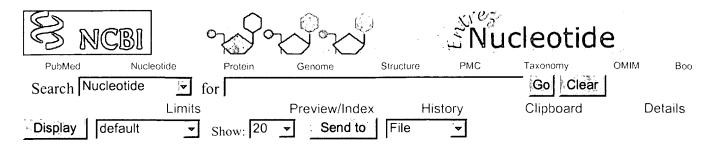
1997

Status:

Unpublished

MAP DATA

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☐ 1: AI041596. ox56h06.x1 Soares...[gi:3280790]

Links

IDENTIFIERS

dbEST Id: 1780931 EST name: ox56h06.x1 GenBank Acc: AI041596 GenBank gi: 3280790

CLONE INFO

Clone Id: IMAGE:1660379 (3')

Source: NCI 827 Insert length: **CDNA** DNA type:

PRIMERS

-40ml3 fwd. ET from Amersham Sequencing:

PolyA Tail: Unknown

SEQUENCE

TGACACCACTGTTGACAATGAAAAAAAAAACAGCATTTGATATTTTCCAGCTTTTTAA GTTAAAAAATGATTCAGTTAAAACAAAACAAAAGTTTAGATATTTTAGCAAGAATCACCT GCAATGTGATTGATTGATTGATTGATATATACCAAAATTCATATGGATGTCACACA

GTCAAGCTAAAAGGAATGTTGGAG

Quality: High quality sequence stops at base: 414

Entry Created: Jun 30 1998 Last Updated: Sep 24 1998

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares_NhHMPu_S1 Homo sapiens Organism: mixed (see below) Organ:

Pooled human melanocyte, fetal heart, and pregnant uterus Tissue type: Lab host:

pT7T3D-Pac (Pharmacia) with a modified polylinker Vector:

R. Site 1: Not I Eco RI R. Site 2:

Description: Equal amounts of plasmid DNA from three normalized libraries

> (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and ss circles were made in vitro.

Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was

PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones

260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

Authors:

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

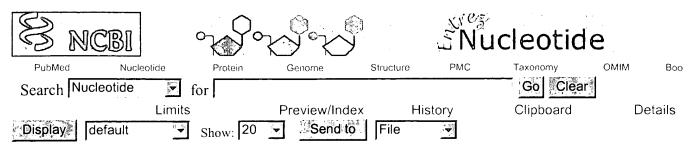
1997

Status:

Unpublished

MAP DATA

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☐ 1: AW072470. xa06d07.x1 Soares...[gi:6027468]

Links

IDENTIFIERS

dbEST Id: EST name: 3209709

GenBank Acc:

xa06d07.x1 AW072470

GenBank gi:

6027468

CLONE INFO

Clone Id:

IMAGE:2567533 (3')

Source: Insert length: NCI 1130

DNA type:

cDNA

PRIMERS

Sequencing:

-40UP from Gibco

PolyA Tail:

Unknown

SEQUENCE

Quality:

High quality sequence stops at base: 447

Entry Created: Last Updated: Oct 13 1999 Oct 20 2000

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium ($\underline{info@image.llnl.gov}$) for further information.

LIBRARY

Lib Name:

Soares_NFL_T_GBC_S1

Organism:

<u>Homo sapiens</u>

Organ: Lab host: pooled

Vooton

DH10B

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I

R. Site 2:

Eco RI

Description:

Equal amounts of plasmid DNA from three normalized libraries (fetal lung NbHL19W, testis NHT, and B-cell NCI_CGAP_GCB1) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive

hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 297480-302087, 682632-687239, 726408-728711, and 729096-731399. Subtraction by Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

Authors:

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

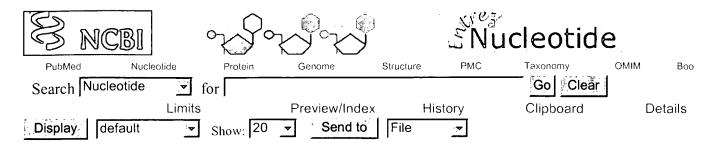
1997

Status:

Unpublished

MAP DATA

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1: AI573107. tr69f01.x1 NCl_CG...[gi:4536481]

Links

IDENTIFIERS

 dbEST Id:
 2391435

 EST name:
 tr69f01.x1

 GenBank Acc:
 AI573107

 GenBank gi:
 4536481

CLONE INFO

Clone Id: IMAGE:2223577 (3')

Source: NCI Insert length: 3468 DNA type: cDNA

PRIMERS

Sequencing: -40UP from Gibco

PolyA Tail: no

SEQUENCE

GGTCAAGCTAAAAGGAATGTTGGAGCCAAAA

Quality: High quality sequence stops at base: 405

Entry Created: Mar 29 1999 Last Updated: May 14 1999

COMMENTS

Life Technologies catalog #: 11548-013

DNA Sequencing by: Washington University Genome Sequencing

Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

LIBRARY

Lib Name: NCI_CGAP_Pan1
Organism: Homo sapiens
Organ: pancreas

Tissue type: adenocarcinoma

Lab host: DH10B

Vector: pCMV-SPORT6

R. Site 1: SalI R. Site 2: NotI Description:

Cloned unidirectionally. Primer: Oligo dT. Average insert

size 1.72 kb. Life Technologies catalog #: 11548-013

SUBMITTER

Name: E-mail: Robert Strausberg, Ph.D. cgapbs-r@mail.nih.gov

CITATIONS

Title:

National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

Authors:

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year:

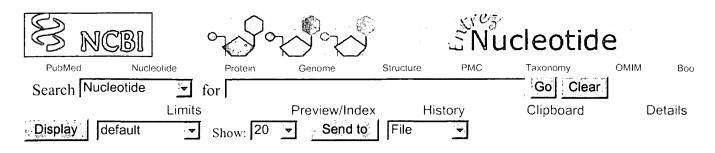
1997

Status:

Unpublished

MAP DATA

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☐ 1: AW770384. hl80d09.x1 NCI_CG...[gi:7702426]

Links

IDENTIFIERS

dbEST Id:

4231435

EST name:

hl80d09.x1

GenBank Acc:

AW770384

GenBank gi:

7702426

CLONE INFO

Clone Id:

IMAGE:3007505 (3')

Source:

NCI

DNA type:

CDNA

PRIMERS

Sequencing:

-40UP from Gibco

PolyA Tail:

Unknown

SEQUENCE

AATGACACCACTGTTGACAATGAAAAAAAAAACAGCATTTGATATTTTCCAGCTTTTT AAGTTAAAAAATGATTCAGTTAAAACAAAACAAAAGTTTAGATATTTTAGCAAGAATCAC ATGCAATGTGATTGATTGATTGATTGATATATACCAAAATTCATATGGATGTCACA TGGTCAAGCTAAAAGGAATGTTGGAGCCAAAAGAAATGT

Quality:

High quality sequence stops at base: 415

Entry Created:

May 4 2000

Last Updated:

May 4 2000

COMMENTS

Tissue Procurement: Chris Moskaluk, M.D., Ph.D., Michael R. Emmert-Buck, M.D., Ph.D. cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: Christa Prange, The I.M.A.G.E. Consortium DNA Sequencing by: Washington

University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL, send

email to: info@image.llnl.gov

LIBRARY

Lib Name:

NCI_CGAP_Kid13

Organism:

Homo sapiens

Organ:

kidney

Tissue type:

2 pooled Wilms' tumors, one primary and one metastatic to

brain DH10B

Lab host: Vector:

pCMV-SPORT6

R. Site 1: SalI R. Site 2: NotI

Description: Cloned unidirectionally. Primer: Oligo dT. Library

constructed by Life Technologies.

SUBMITTER

Name: Robert Strausberg, Ph.D. E-mail: cgapbs-r@mail.nih.gov

CITATIONS

Title: National Cancer Institute, Cancer Genome Anatomy Project

(CGAP), Tumor Gene Index

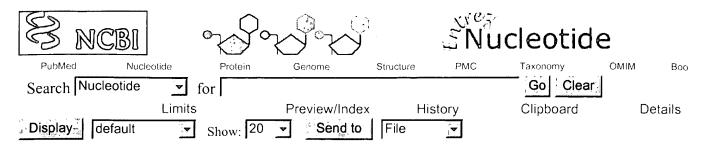
Authors: NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap

Year: 1997

Status: Unpublished

MAP DATA

<u>Disclaimer</u> | <u>Write to the Help Desk</u> <u>NCBI</u> | <u>NLM</u> | <u>NIH</u>



☐ 1: AA479302. zv21f08.s1 Soares...[gi:2207858]

Links

IDENTIFIERS

 dbEST Id:
 1124337

 EST name:
 zv21f08.s1

 GenBank Acc:
 AA479302

 GenBank gi:
 2207858

 GDB Id:
 5977197

CLONE INFO

Clone Id: IMAGE: 754311 (3')

Source: IMAGE Consortium, LLNL

DNA type: cDNA

PRIMERS

Sequencing: -41m13 fwd. ET from Amersham

PolyA Tail: Unknown

SEQUENCE

AAGCTAAAAGGAATGTTGGAGCCAAAAGAAATGTAGTGGAAT

Quality: High quality sequence stops at base: 456

Entry Created: Jun 19 1997 Last Updated: Aug 8 1997

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares_NhHMPu_S1
Organism: Homo sapiens
Organ: mixed (see below)

Tissue type: Pooled human melanocyte, fetal heart, and pregnant uterus

Lab host: DH10B

Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI

Description: Equal amounts of plasmid DNA from three normalized libraries

(melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and ss circles were made in vitro.

Following HAP purification, this DNA was used as tracer in a

subtractive hybridization reaction. The driver was

PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones 260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

Name: Wils

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel:

314 286 1800 314 286 1810

Fax: E-mail:

est@watson.wustl.edu

CITATIONS

Title:

WashU-Merck EST Project 1997

Authors:

Hillier,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising

,B., White,Y., Wylie,T., Waterston,R., Wilson,R.

Year:

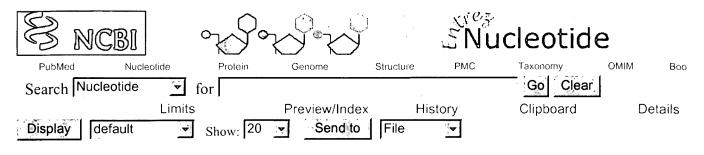
1997

Status:

Unpublished

MAP DATA

Disclaimer | Write to the Help Desk NCBI | NLM | NIH



1: R27957. yh57b02.s1 Soares...[gi:784092]

Links

IDENTIFIERS

 dbEST Id:
 198002

 EST name:
 yh57b02.s1

 GenBank Acc:
 R27957

 GenBank gi:
 784092

 GDB Id:
 539557

CLONE INFO

Clone Id: IMAGE:133803 (3')

Insert length: 820 DNA type: cDNA

PRIMERS

Sequencing: Promega -21m13

PolyA Tail: Unknown

SEQUENCE

NATACTTGGGGNATGGAGGGCTGTGG

Quality: High quality sequence stops at base: 302

Entry Created: Apr 25 1995 Last Updated: Apr 25 1995

COMMENTS

Insert Size: 820

High quality sequence stops: 302 Source: IMAGE Consortium, LLNL

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (<u>info@image.llnl.gov</u>) for further

information.

LIBRARY

Lib Name: Soares placenta Nb2HP

Organism: Homo sapiens

Sex: Female Organ: placenta

Develop. stage: placenta obtained at birth (full term)

Lab host: DH10B (ampicillin resistant)

Vector: pT7T3D (Pharmacia) with a modified polylinker

R. Site 1: Not I R. Site 2: Eco RI Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

double-stranded cDNA was ligated to Eco RI adaptors

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by

Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

Wilson, R.

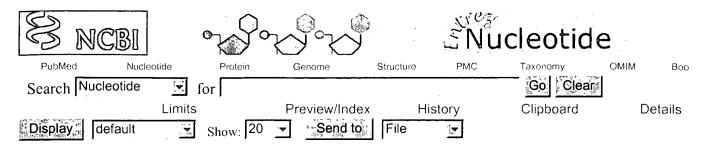
Year: 1995

Status: Unpublished

MAP DATA

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Jul 8 2003 12:22:35



☐ 1: H05625. yl70b12.s1 Soares...[gi:869177]

Links

IDENTIFIERS

 dbEST Id:
 265505

 EST name:
 y170b12.s1

 GenBank Acc:
 H05625

 GenBank gi:
 869177

 GDB Id:
 415859

CLONE INFO

Clone Id: IMAGE:43318 (3')

DNA type: cDNA

PRIMERS

Sequencing: Promega -21m13

PolyA Tail: Unknown

SEQUENCE

ACCGTAATCTGGGCAACATCCGGGGGCTTACCTTCAGCTCTCGGCACTGTGCG

Quality: High quality sequence stops at base: 310

Entry Created: Jun 21 1995 Last Updated: Jun 21 1995

COMMENTS

High quality sequence stops: 291 Source: IMAGE Consortium, LLNL

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name: Soares infant brain 1NIB

Organism: Homo sapiens
Sex: female
Organ: whole brain

Develop. stage: 73 days post natal

Lab host: DH10B (ampicillin resistant)

Vector: Lafmid BA
R. Site 1: Not I
R. Site 2: Hind III

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

(Pharmacia), digested with Not I and directionally cloned into the Not I and Hind III sites of the Lafmid BA vector. Library went through one round of normalization. Library constructed by Bento Soares and M.Fatima Bonaldo.

SUBMITTER

Name: Wilson RK

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Tel: 314 286 1800 Fax: 314 286 1810

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CITATIONS

Title: The WashU-Merck EST Project

Authors: Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,

Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M., Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

Wilson, R.

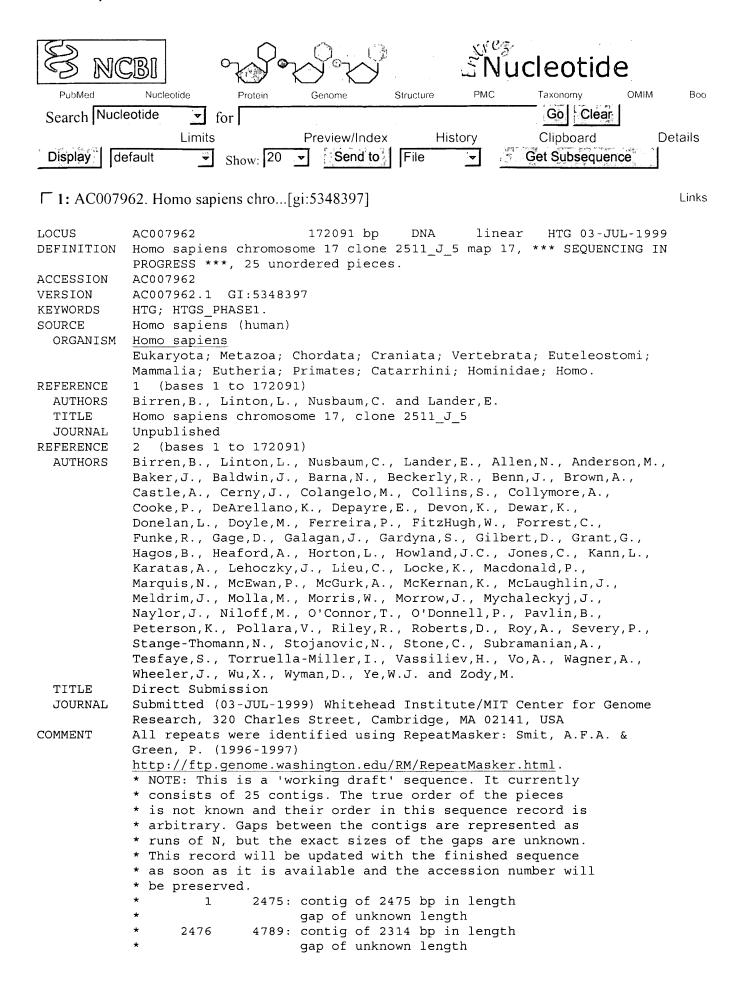
Year: 1995

Status: Unpublished

MAP DATA

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Jul 8 2003 12:22:35



4790

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7000: contig of 2211 bp in length

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